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# Recombinant full-length tissue factor pathway inhibitor fails to bind to the cell surface: implications for catabolism in vitro and in vivo

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Tissue factor pathway inhibitor (TFPI) plays a key role in the regulation of tissue factor-initiated blood coagulation secondary to loss of the integrity of the blood vessel wall. TFPI is a naturally occurring Kumitz-type protease inhibitor that inhibits coagulation factor Xa and, in a factor Xa-dependent manner, mediates feedback inhibition of the factor Vilafitssuefactor

catalytic complex. In vivo full-length TFPI is thought to be primarily bound to the vascular endothelium and the high affinity binding requires an intact carboxy terminus. Here we describe a full-length TFPI molecule, expressed in mouse C127 cells (TFPI<sup>C127</sup>), which exhibits virtually no cellular binding yet contains the intact carboxy terminus. This TFPI (TFPI<sup>C127</sup>) is

neither internalized nor degraded via the TFPI endocytic receptor, LDL-receptor-related protein. Pharmacoldnetic studies of TFPI<sup>CED</sup> in vivo demonstrate a 10-fold prolongation in the plasma half-life, compared with that of bactarial recombinant TFPI. (Blood, 2000;95:1973-1978)

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# Introduction

The extrinsic pathway of coagulation is initiated by the exposure of tissue factor (TF) to the circulating blood. Binding of factor VII to TF promotes the activation of factor VII to factor VIIa. The factor VIIa/TF complex activates its substrates factor IX and factor X, triggering a cascade of events that culminates in the formation of a fibrin clot.<sup>23</sup> Tissue factor pathway inhibitor (TFPI), a 42 kd plasma glycoprotein, plays a key role in the regulation of TFinitiated blood coagulation23 via its abilities to directly inhibit factor Xa and, in a factor Xa dependent manner, inhibit the factor VIIa-TF proceedytic activity. Mature TFPI consists of a negatively charged amino terminus, 3 tandom Kunitz-type inhibitory domains, and a positively charged carboxy terminus. The second Kunitz domain binds and inhibits factor Xa, while the first Kunitz domain binds and inhibits the factor VIIa/TF complex.4 The carboxy terminus of TFPI and a portion of the third Kunitz domain contain heparin-binding sites 16 and are necessary for the anticoegulant function of TFPI in TF-induced coagulation in vitro<sup>5</sup> and for TFPI binding to the cell surface.7

TFPI circulates in plasma predominantly bound to lipoproteins. Platelets contain approximately 8% of the total TFPI in blood and release their TFPI after stimulation with thrombin. The major endogenous source of TFPI is thought to be bound to the vascular endothelium and is releasable after heparin infusion, whereupon plasma TFPI levels rise several fold. 10,11

Recent animal studies have demonstrated that recombinant TFPI is effective against TF-induced coagulopathy, <sup>12</sup> prevents arterial thrombosis, <sup>13,14</sup> and reduces mortality from bacterial septic shock. <sup>15</sup> Pharmacokinetic studies after an intravenous bolus injection of recombinant TFPI have shown that TFPI is rapidly cleared from the circulation with a plasma half-life of approximately 2 minutes in rabbits <sup>16</sup> and less than 1 minute in rate. <sup>7</sup> As a result, high doses of recombinant TFPI (approximately 20 mg/kg/d) are required to echieve therapeutic efficacy. <sup>17</sup>

We have recently shown that the rapid clearance of recombinant TFPI from the circulation is in large part dependent on the binding of TFPI to the vascular endothelium and to a lesser degree to its hepatic removal via the endocytic receptor LDL-receptor-related protein (LRP). In this study, we demonstrate that full-length TFPI expressed in mouse C127 cells does not bind to the cell surface nor is it degraded via LRP in vitro. This results in an approximately 10-fold prolongation of the TFPIC127 plasma half-life in mice. The noncellular binding properties of TFPIC127 may thus offer additional therapeutic potential.

# Materials and methods

## Materials

lodogen was purchased from Pierce (Reckville, IL). [137] Ilodida was from Amersham Corp (Pierataway, NJ). Human factor Xa was from Enzyme Research (South Bend, IN) and Spectruzyme Xa was from American Diagnostica, Inc (Greenwich, CT). Bovine serven ablumia was purchased from Collection-Novabiochem. Affi-gel 10 was from Bio-rad (Hercules, CA). Tissun culture modia and plastic-ware were obtained from Life Technologies, Inc (Rockville, MD). Recombinant TFPI from Encherichia coli was provided by Monsanto Company (St Louiz, MO). CHO, SK hepstoma, and 293 cell-derived TFPI, was purified as described below.

Parification of TFP1<sup>CLS7</sup>—TFP1<sup>CLS7</sup> was affinity-purified from the conditioned media of mouse C127 cells that had been transfected with a bovine papilloma virus containing wild-type human TFP1 cDNA. <sup>12</sup> In brief, cells producing TFP1<sup>CLS7</sup> were cultured in serum-free media containing apprainin (10 µg/mL; Sigma) and the media were replaced every 2 to 3 days. Harvested media were concentrated using an Amicone observator (YM 30; Amicon, Bedford, MA) and the concentrates were absorbed to a 2H8 monocloual anti-TFP1-affigel 10 column (0.5 × 3 cm) and chief with 100 mmol/L glycine, pH 2.2. The pH of the chutant was adjusted to pH 7.5 using 1 mol/L Tris-HCl, pH 8.0, and applied to a 1 mL HiTrap

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heparin-agarose column (Pharmacia-LKB Biotechnology, Piscataway, NJ), which had been equilibrated in 0.1 mol/L NaCl, 0.05 mol/L Tria-HCl, pH 7.5. The column was developed with a linear NaCl gradient from 0.1 to 1.0 mol, at 0.5 ml/min over 10 column volumes. Practions were assayed for the presence of TFPICET by a modified enzyme linking immunosorbent assay (RLISA) method using 2 independent monoclonal antibodies, 2H8 and 2B12. The desired fractions were pooled, concentrated using Centricon microconcentrators (Amicon), and stored at ~80°C. The protein concentration was determined using the modified ELISA method with TFPI of known concentration as attendard.

## Amidolytic assays of factor Xa activity

In a 96-well microtier plain. 1 mmsl/L of factor Xa was incubated with various concentrations of TFPI (0-3 mmsl/L) in 135 pL of 0.1 mol/L NaCl, 0.05 mol/L Pris-HCl, 0.5% bovine serum abbumin (BSA), pH 7.5 (TBSA) at room temperature for 30 minutes. Fifteen microtiters of 1 mmsl/L Spectrozyme Xa chromagenic substrate was then introduced (final concentration 100 pmsl/L) and the change in absorbance at 405 mm was determined 10 minutes thereafter.

# Tissue factor pathway inhibitor functional assay of Vila/TF inhibition

The end point assay of anti-VIIA/TF activity (American Diagnostica) was used to compare TFPf<sup>CE27</sup> and bacterially expressed TFPI at equimolar concentrations.

## Factor Xa-induced coagulation of plasma

in a fibrometer (BBL, Cockeysville, MD), 50 µL of rabbit brain cephalin prepared as described by the manufacturer (Signas), 50 µL of CaCl<sub>2</sub> (25 mmol/L), 50 µL of various amounts of TPPI<sup>CL2</sup> (25 ng to 300 ng) in TBSA, and 50 µL of human factor Xa (0.2 nmol/L) were incubated at 37°C. After 30 seconds, 50 µL of normal human plasma (George King Blochemical) were added, and the degree of apparent factor Xa inhibition was determined by comparing the clotting to a factor Xa standard curve.

## Call culture

Human hepatoma Hep (12 cells, mouse fibroblast PBA 13 cells, <sup>15</sup> and mouse bin endothelial cells (bend-3)<sup>20</sup> were enlimed in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% fetal culf scrum, 100 U/mL of penicillin, and 100 µg/mL of streptomycin. Cells were incubated at 37°C in humidified air containing 5% CO<sub>2</sub>.

# Protein logination

Proteins (5-25 µg) were indimated using the IODOGEN method. 21 Specific radioscrivings were typically 1 to 10 × 10 com/pmol of protein.

# Binding and degradation assays

Binding of <sup>125</sup>I-TFPI to cells was performed in suspension. Cells were displaced from Petri-dishes by incubation with 3 mmol/L EDTA in phosphate-buffered seline (PBS) at room temperature for 5 minutes. Cells were then pelleted at low speed (1000 ppm × 5 minutes) on a table top cintrifuge (Sorvall R76000B) to remove EDTA and resuspended in the assay buffer (DMEM/3% BSA). 4 × 10<sup>5</sup> cells in a volume of 300 µL were added with 2 mol/L of <sup>125</sup>I-TFPI in the presence or absence of 100 mol excess of cold ligands. After rocking at 4°C for 2 hours, cell suspensions were gently toyered on 900 µL of first cell serious in microfuge tabes and spun at 14 000 rpm × 1 minute in a table-top microfuge. The cell pellet was counted in a gamma counter (Cobra II sub-gamma, Packard Instrument, Meredin, CT). Nonspecific binding was determined in the presence of circus unlabeled ligand as specified in the figure legends.

Degradation assays were purformed by washing cell monolayers (in 12-well disters) twice with the assay buffer. 0.5 mL of assay buffer containing 2 nmol/L <sup>12</sup>L-TFPI was then added to each well. After incubation at 37°C for the indicated periods, the overlying medium was removed and proming were procephaned by the addition of BSA to 5 mg/ml.

and trichloroscetic soid to 20%. Degradation of ligand was defined as the appearance of radioactive fragments in the overlying medium that were soluble in trichloroscetic acid. Pragmentation of <sup>127</sup>1-TFPI at 37°C in cell-free wells was substructed from the corresponding samples.

For ligand competition binding and degradation assays (Table 2), HepO2 and CEO cells were assayed as monolayers as described previously<sup>7,16</sup>, <sup>125</sup>1-TFPI was at 20 nmol/L for the binding assays and at 0.6 pmol/L for the degradation assays.

## Plasma clearance of <sup>528</sup> (issue factor pathway inhibitors in mice

Twelve- in 16-week-old BALB/e mice (weighing 20-25 g) were sneatherized with sodium pentobarbinal (1 mg/20 g of body weight) during the course of experiments. Approximately 15 pmol of radiolabeled protein slone or with unkneled 39 lnd protein in 100 µL of sterile saline was injected into the tail vein over 30 seconds. In studies in which heparin was administered, heparin was injected 20 minutes after <sup>125</sup>-TFPI administration. Al the indicated times, 40 to 50 µL of blood was collected by periorbinal bleeding. The blood samples were centrifuged and 15 µL of the plasma fractions was spotted onto 3 MM filter paper (Whatman Loe, Clifton, NI), precipitated with 10% trichloroscetic acid, rinsed with ethanol, and radioactivity determined. The initial plasma concentration (time = 0) of <sup>123</sup>-TPPI was extrapolated from the measured points. At the and of each experiment, unfinals were killed and the liver, kidneys, spleen, and lungs were removed, blotted, weighed, and the radioactivity determined.

# Metabolic incorporation of <sup>36</sup>SO<sub>4</sub> and immunoprecipitation of TEPI<sup>C127</sup>

One 75-cm<sup>2</sup> flask of C127 cells was labeled for 96 hours with 37 MBq (I mCi) of Na<sub>2</sub>[<sup>25</sup>S]SO<sup>4</sup> (Amersham Corp) in 10 mL of sulfain-deficient DMEM (-SO<sub>4</sub>) supplemented with 2% dialyzed fetal calf serum, in the absence of G418. The culture medium was then concentrated using Centricon microconcentrators (Amicon). Monoclonal antihody 2H8-conjugated affigel-10 beads were added to the concentrates that were then rocked overnight in allow for binding. The beads were subsequently collected by centrifugation, washed 3 times with PBS, boiled in SDS-sample buffer (to release bound TFPI), and the bead-free portion was analyzed by SDS-PAGE.

# Electrophorasis and Western blotting

SDS-PAGE was performed using 10% separating and 5% stacking gets and proteins were transferred to nitrocellulose membrane. The membrane was blocked with 5% nonfax milk for 30 minutes, followed by incubation with a 1:100 dilution of rabbit antisera raised against a synthetic peptide matching the carboxy terminal 12 amino acid residues of the manuse TPPI protein. 

The incompresentivity was detected using BCL reagents (Ameraham Corp).

# Results

# Purification of recombinant tissue factor pathway inhibitor from mouse C127 cells

Previous studies have shown that TFPI, when expressed in mammalian systems, is susceptible to protoolysis at its carboxy terminus. TFPI was grown in media commining the protease inhibitor, aprotinin. TFPI was purified from the media via TFPI monoclonal antibody-affinity chromatography. This was followed by heparla-agarose affinity which separates the full-length TFPI from the truncated species by charge interactions between the heperin and the positively charged carboxy terminus of TFPI. Figure 1 shows the clution of TFPICE27 from the heparin column with a linear NaCl gradient. TFPICE27

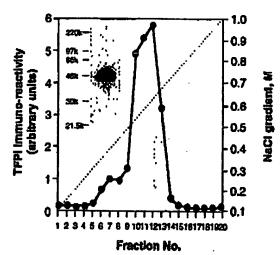


Figure 1. Profile of Inspirits-egators chromategraphy of TEP(\*\*12\*), Hopprinagerose chromatography was carried out using a 0.1 to 1 molf. NGCI gradient in 0.05 molf. Tris-HOI, pH 7.5. Elution of TEP(\*\*22\*) was monitored by a modified ELISA as described in the "Haterials and Mothads." Inset: Western blot analysis of TEP(\*\*2\*\*). Purified TEP(\*\*2\*\*) (1 pg) was non on a 10% SDS-PAGE gail and liansterned to introcebulose membrane. The membrane was blotted with rabbit andexes containing a people antibody against the last 12 amino acid residues of TEPI, and developed with ECI, regions.

etuted with 2 peaks at NaCl concentrations of 0.4 and 0.6 mol/L, respectively. Because full-length TFP1 clutes at 0.6 mol/L salt. 5.6 only the fractions collected at or greater than 0.6 mol/L NaCl (fractions 12-15) were pooled for subsequent studies. To confirm that the purified TFPICIT was full length, TFPICIT was subjected to Western blot analysis using antibodies directed against the last 12 amino acid residues of TFPI. As shown in the inset of Figure 1, the TFPI was recognized by these antibodies, confirming that the carboxy terminus was intact.

The activity of TFPICLY was assessed in a direct amidolytic assay of factor Xa activity, as well as an end point assay of

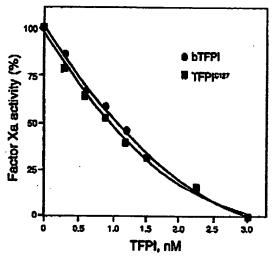


Figure 2. Inhibition of factor Xa amidolytic activity by TEPIFED. Factor Xb (1 mmd) was incubated with increasing concentrations of TEPIFED (0-3 mmd) at room temporature for 30 minutes. The remaining factor Xa activity was assessed using the chromagonic substrate Spectrosymb Xa. Backets recombined TEPI (bTFPI) was assessed in the Identical minuter and used as control.

Table 1. Blodleg of VIALTEPICIET to vertous cell knee

Cell Lires	Specific Binding (Imal/10 <sup>a</sup> calls)	
	1314-Athle152	C21-6TFPI
HopG2	Not detectable	590
Bond	Not detectable	810
PEA 13	Not detectable	540

HapG2, Bend, and PEA 19 cells were incubated in suspension with 2 most traintenance or absence of 100 mol excess of cold TEPICTY in the presence or absence of 100 mol excess of cold TEPICTY for 2 hours at 4°C. The specific redisactivity sesociated with colls was determined (open bard). Binding of bacterial recombinant <sup>123</sup>-TEPI to these calls was performed in the fearful meaner and used as control. Values represent the means of duplicate decombinations.

anti-VIIA/TF activity. As shown in Figure 2, TFPI<sup>C127</sup> was able to inhibit factor Xa activity in a concentration-dependent manner with complete inhibition of factor Xa at approximately 3 mmol/L, which was essentially identical to that of bacterial recombinent TFPI Using a global anticoagulant assay, TFPI<sup>C127</sup> inhibited factor Xa-induced plasma coagulation with IC<sub>30</sub> approximately 10 nmol/L, compared with approximately 4 mmol/L for the bacterial recombinant TFPI. The disparity in IC<sub>30</sub> values observed here is consistent with the previous observations that TFPI expressed in prokaryotic systems exhibits higher anticoagulant activities than that expressed in cultaryotic systems. Using the end pour assay that measures anti-VIIA/TF activity, on a molar the activity of TFPI<sup>C121</sup> is 108% ± 16% that of bacterially expressed TFPI.

# Cellular binding and degradation of tissue factor pathway inhibitor in C127 cells

We have previously shown that bacterial recombinant TFPI binds to human hepatoma HcpG2 cells with high affinity<sup>34</sup> and that truncation of sequences distal to the second Kunitz domain abolishes the binding.<sup>7</sup> Surprisingly, the full-length <sup>123</sup>I-TFPiC<sup>123</sup> exhibited no binding to hepatoma HcpG2 cells, or to mouse vascular endothelial Bend cells, or mouse fibroblast PEA 13 cells (Table 1). Bacterial recombinant <sup>125</sup>I-TFPI, however, exhibited abundant binding to these cells (Table 1). Furthermore, <sup>125</sup>I-TFPIC<sup>127</sup> was not taken up/degraded by HcpG2 cells in contrast to

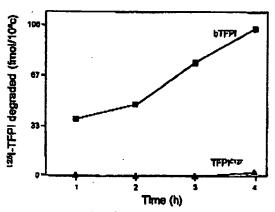


Figure 3. Degredation of <sup>128</sup>L-TFPI<sup>C-97</sup> by HapG2 calls. HapG2 palls growing in potel dishes were incubated with 0.6 mmol of <sup>128</sup>L-TFPI<sup>C-97</sup> at 37°C for the indicated limes. At each time point, the overlying medium was removed and proteins were precipitated by Withhospeeds and (TCA). The TCA-totate redonately (is. degression products) was determined. Degradation of bacterial recombinest <sup>128</sup>L-TFPI by HapG2 calls was performed in the Identical manner and served as a certaria. Numbers represent the means of duplicate determinations. Total degraded <sup>128</sup>L-TFPI at 4 hours was less than 10% hours.

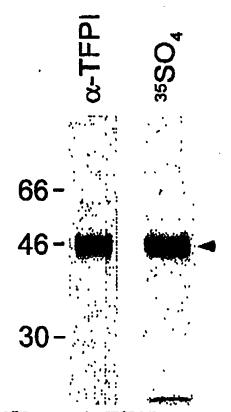


Figure 5. MSO<sub>4</sub> is incorporated into TFPP<sup>DCD</sup>, C127 cells expressing human TFPI were included with <sup>25</sup>CO<sub>4</sub> for 85 hours, after which the culture medium was collected and immunoprecipitated with anti-TFP as described under "Naturals and Mediuds," The immunoprecipitates were analyzed vie 805-PACE and extendiography (right), or via Wastern biot analysis using TFPI monocloset enthody 2H8 (ods).

Binding of full-length TFPI to the cell surface is dependent on its highly positively charged carboxyl terminus that interacts with the negatively charged cell surface HSPGs.7,10,28 We have proviously shown that TFPI expressed in bacteria binds with high affinity to heparoma HepG2 cells, fibroblast PEA13 cells, and microvascular endothelial cells. 20,24,27 Similarly, TPPI expressed in hopatoma and CHO cells displays similar binding properties (Table 3). To our surprise, TFPI expressed in mouse C127 fibroblest cells was devoid of cell surface binding activity but remained biologically active as judged by the inhibition of (1) factor Xa amidolytic activity (Figure 2), of (2) factor Xa-induced plasma congulation, and (3) factor VIIs/TF activity. One potential explanation for this loss of binding activity is that secondary to intramolecular intersetions, TFPI's carboxyl terminus is no longer available for interaction with cell surface HSPGs. It is of note that, although TPPICLE does not blind to cell surface HSPGs, it binds to heparin-agarose (Figure 1) with the same affinity as becterial recombinant TFPI, is, both TFPI species are cluted from heparin-agarose at identical salt concentrations (0.6 mol/L NaCl) (6, data not shown). The fact that the affinity of TFPI is 40-fold greater for heparin than for HSPGs28 may account for the high affinity binding of both species of TFPI to heasrin and their discordant binding to cell surface HSPGs.

Girard et al<sup>29</sup> have shown that serine 2 of TFPI is phosphorylated after its expression in C127 cells. However, TFPI with a serine 2 to alanime motation, when expressed in these cells, did not regain the binding properties of wild type TFPI (data not shown). This finding suggests that this negatively charged phosphate group (ie, at serine 2) does not interfere with availability of the carboayl terminus. TFPI contains 3 potential N-linked and 1 O-linked glycosylation sites. 30 Smith et al 24 have shown that metabolic labeling of 293 cells with [35]SO4 yielded sulfated TFPI, whereas sulfated TFPI was not produced by CHO cells. They further determined that TFPi<sup>29)</sup> carries Asn-linked sulfaced oligosaccharides: SO\_GaINAc61,2GlcNAc61,2Mana.25 Because TFP1297 was unable to bind to the cell surface (Table 3), it is possible that sulfated oligosaccharides on TPPI may interact with its own positively charged carboxyl terminus, thus abeogating TFPI's interaction with the cell surface. Attempts to quantitatively remove linked oligosaccharides or sulfate via incubation of nondenanced TFPIC127 with N-glycenese, O-glycenese or sulfanse and assess biological function have proven unsuccessful. Thus, whether TFPI<sup>C122</sup> possesses the same sulfation pattern as TFPI<sup>233</sup> will require further study.

TFPICIZI exhibited a 10-fold increase in plasma half-life compared with bacterial recombinant TFPI (Figure 4A). Because the administration of heparin did not significantly increase the plasma <sup>125</sup>I-TFPI<sup>CLD</sup> concentration, secondary to release from the vascular endothellum, or did the administration of the 39-kd protein inhibit its clearance (Figure 4), the prolonged half-life of FIFE-TFPICITY is likely due to its inability to bind to either cell surface HSPGs or LRP.31 These results are consistent with our in vitro findings (Figure 3 and Table 1). However, it is not clear at present whether this observation can be accounted for by alterations in procein sulfation alone, because 1251-TFP1293 and 1251-TFPICHO have similar rates of plasma clearance.26 Nonetheless, the clearance of 1251-TFPIC127, albeit markedly prolonged, did result in elimination of 50% at approximately 20 minutes. Therefore, additional mechanisms (eg. hepatic glycoprotein receptors) may be operative.

Under physiologic conditions, TFPI is primarily synthesized and secreted by vascular endothelial cells. Because the major pool of TFPI in vivo is associated with the endothelial cell surface, it appears that endothelial TFPI maintains the carboxy terminus available for inverserion with cell surface HSPGs. Assuming that sulfation does play a role in TFPI binding to cells, TFPI produced by endothelial cells in vivo may well have to be under- or unsulfated to maintain its cellular binding. However, it is not presently known whether endogenous TFPI is sulfated or its degree of sulfation. Of note, human endothelial cells express significantly lower (less than 10%) GalNAc- and sulforansferase activities than those expressed in 293 cells. These 2 enzymes are the key activities required for the addition of sulfated oligosaccharides to TFPI.26

The physiologic significance of the cell-associated TFPI pool may relate to both its rapid mobilization into the plasma as well as its role as a mediator in the cellular uptake and degradation of factor Xa.<sup>27</sup> In addition, it may serve to down-regulate the coagulant activity of factor VIIa/TF complex via transfer to cavoolae.<sup>25</sup> Endogenous TFPI within the circulation, however, is likely of less physiological importance, as it is largely truncated at various positions within the carboxy terminal domain<sup>26</sup> and hence exhibits much reduced anticoagulant activity.<sup>3</sup> The nonbinding full-length TFPI we have characterized in this study may therefore offer a therapeutic advantage when high plasma TFPI concentrations are required to effectively modulate TF-induced blood coagulation.

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